

SEQUENCE LISTING

ŽEUŢLER, BRUCE POLTORAK, ALEXANDER

<120> LPS - RESPONSE GENE COMPOSITIONS AND METHODS

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<170> PatentIn Ver. 2.1

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Ile Tyr Asp Ala Phe Val Ile Tyr Ser Ser Gln Asp Glu Asp Trp Val 675 680 685

Arg Asn Glu Leu Val Lys Asn Leu Glu Glu Gly Val Pro Pro Phe Gln 690 695 700

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- Thr Phe Pro Tyr Lys Cys Leu Asn Ser Leu Gln Val Leu Asp Tyr Ser
- Leu Asn His Ile Met Thr Ser Lys Gln Glu Leu Gln His Phe Pro
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- Gly Met Pro Val Leu Ser Leu Asn Ile Thr Cys Gln Met Asn Lys Thr
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- Val Leu Val Tyr Lys Phe Tyr Phe His Leu Met Leu Leu Ala Gly Cys
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- Ser Ser Gln Asp Glu Asp Trp Val Arg Asn Glu Leu Val Lys Asn Leu
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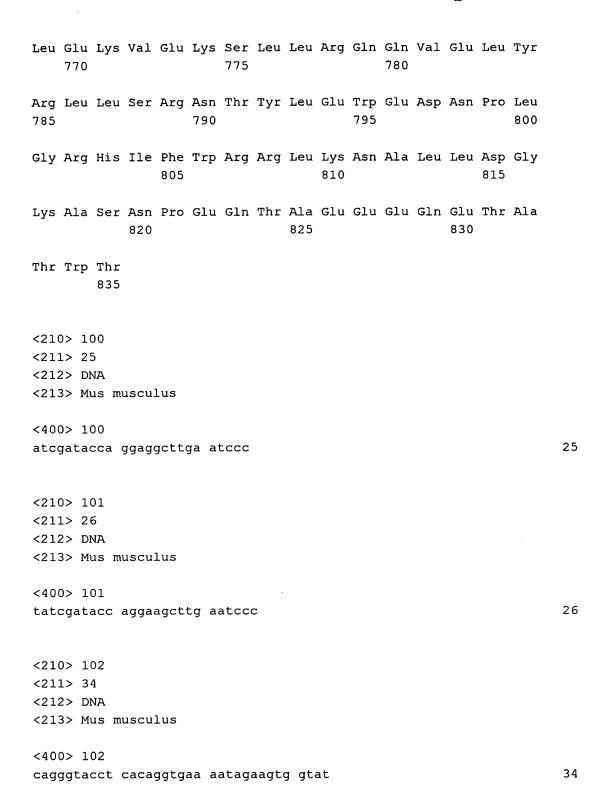
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Tyr	Leu 450	Asp	Ile	Ser	Tyr	Thr 455	Asn	Thr	Lys	Ile	Asp 460	Phe	Asp	Gly	Ile
Phe 465	Leu	Gly	Leu	Thr	Ser 470	Leu	Asn	Thr	Leu	Lys 475	Met	Ala	Gly	Asn	Ser 480
Phe	Lys	Asp	Asn	Thr 485	Leu	Ser	Asn	Val	Phe 490	Ala	Asn	Thr	Thr	Asn 495	Leu
Thr	Phe	Leu	Asp 500	Leu	Ser	Lys	Cys	Gln 505	Leu	Glu	Gln	Ile	Ser 510	Trp	Gly

- Val Phe Asp Thr Leu His Arg Leu Gln Leu Leu Asn Met Ser His Asn 515 520 525
- Asn Leu Leu Phe Leu Asp Ser Ser His Tyr Asn Gln Leu Tyr Ser Leu 530 540
- Ser Thr Leu Asp Cys Ser Phe Asn Arg Ile Glu Thr Ser Lys Gly Ile 545 550 555 560
- Leu Gln His Phe Pro Lys Ser Leu Ala Phe Phe Asn Leu Thr Asn Asn 565 570 575
- Ser Val Ala Cys Ile Cys Glu His Gln Lys Phe Leu Gln Trp Val Lys 580 585 590
- Glu Gln Lys Gln Phe Leu Val Asn Val Glu Gln Met Thr Cys Ala Thr
 595 600 605
- Pro Val Glu Met Asn Thr Ser Leu Val Leu Asp Phe Asn Asn Ser Thr 610 620
- Cys Tyr Met Tyr Lys Thr Ile Ile Ser Val Ser Val Val Ser Val Ile 625 630 635 640
- Val Val Ser Thr Val Ala Phe Leu Ile Tyr His Phe Tyr Phe His Leu 645 650 655
- Ile Leu Ile Ala Gly Cys Lys Lys Tyr Ser Arg Gly Glu Ser Ile Tyr
 660 665 670
- Asp Ala Phe Val Ile Tyr Ser Ser Gln Asn Glu Asp Trp Val Arg Asn 675 680 685
- Glu Leu Val Lys Asn Leu Glu Glu Gly Val Pro Arg Phe His Leu Cys 690 695 700
- Leu His Tyr Arg Asp Phe Ile Pro Gly Val Ala Ile Ala Ala Asn Ile 705 710 715 720
- Ile Gln Glu Gly Phe His Lys Ser Arg Lys Val Ile Val Val Ser 725 730 735
- Arg His Phe Ile Gln Ser Arg Trp Cys Ile Phe Glu Tyr Glu Ile Ala
 740 745 750
- Gln Thr Trp Gln Phe Leu Ser Ser Arg Ser Gly Ile Ile Phe Ile Val 755 760 765



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Phe Ser Cys Leu Thr Pro Gly Ser Leu Asn Pro Cys Ile Glu Val Val 20 25 30

Pro Asn Ile Thr Tyr Gln Cys Met Asp Gln Lys Leu Ser Lys Val Pro 35 40 45

Asp Asp Ile Pro Ser Ser Thr Lys Asn Ile Asp Leu Ser Phe Asn Pro 50 55 60

Leu Lys Ile Leu Lys Ser Tyr Ser Phe Ser Asn Phe Ser Glu Leu Gln 65 70 75 80

Trp Leu Asp Leu Ser Arg Cys Glu Ile Glu Thr Ile Glu Asp Lys Ala 85 90 95

Trp His Gly Leu His His Leu Ser Asn Leu Ile Leu Thr Gly Asn Pro 100 105 110

Ile Gln Ser Phe Ser Pro Gly Ser Phe Ser Gly Leu Thr Ser Leu Glu
115 120 125

Asn Leu Val Ala Val Glu Thr Lys Leu Ala Ser Leu Glu Ser Phe Pro 130 135 140

Ile Gly Gln Leu Ile Thr Leu Lys Lys Leu Asn Val Ala His Asn Phe 145 150 155 160

Ile His Ser Cys Lys Leu Pro Ala Tyr Phe Ser Asn Leu Thr Asn Leu 165 170 175

Val His Val Asp Leu Ser Tyr Asn Tyr Ile Gln Thr Ile Thr Val Asn 180 185 190

Asp Leu Gln Phe Leu Arg Glu Asn Pro Gln Val Asn Leu Ser Leu Asp 195 200 205

- Ile Met Lys Thr Cys Leu Gln Asn Leu Ala Gly Leu His Val His Arg \$245\$ \$250\$ \$255\$
- Leu Ile Leu Gly Glu Phe Lys Asp Glu Arg Asn Leu Glu Ile Phe Glu 260 265 270
- Pro Ser Ile Met Glu Gly Leu Cys Asp Val Thr Ile Asp Glu Phe Arg 275 280 285
- Leu Thr Tyr Thr Asn Asp Phe Ser Asp Asp Ile Val Lys Phe His Cys 290 295 300
- Leu Ala Asn Val Ser Ala Met Ser Leu Ala Gly Val Ser Ile Lys Tyr 305 310 315 320
- Leu Glu Asp Val Pro Lys His Phe Lys Trp Gln Ser Leu Ser Ile Ile 325 330 335
- Arg Cys Gln Leu Lys Gln Phe Pro Thr Leu Asp Leu Pro Phe Leu Lys 340 345 350
- Ser Leu Thr Leu Thr Met Asn Lys Gly Ser Ile Ser Phe Lys Lys Val 355 360 365
- Ala Leu Pro Ser Leu Ser Tyr Leu Asp Leu Ser Arg Asn Ala Leu Ser 370 380
- Phe Ser Gly Cys Cys Ser Tyr Ser Asp Leu Gly Thr Asn Ser Leu Arg 385 390 395 400
- His Leu Asp Leu Ser Phe Asn Gly Ala Ile Ile Met Ser Ala Asn Phe
 405 410 415
- Met Gly Leu Glu Glu Leu Gln His Leu Asp Phe Gln His Ser Thr Leu 420 425 430
- Lys Arg Val Thr Glu Phe Ser Ala Phe Leu Ser Leu Glu Lys Leu Leu 435 440 445
- Tyr Leu Asp Ile Ser Tyr Thr Asn Thr Lys Ile Asp Phe Asp Gly Ile 450 455 460

Phe Leu Gly Leu Thr Ser Leu Asn Thr Leu Lys Met Ala Gly Asn Ser Phe Lys Asp Asn Thr Leu Ser Asn Val Phe Ala Asn Thr Thr Asn Leu Thr Phe Leu Asp Leu Ser Lys Cys Gln Leu Glu Gln Ile Ser Trp Gly Val Phe Asp Thr Leu His Arg Leu Gln Leu Leu Asn Met Ser His Asn Asn Leu Leu Phe Leu Asp Ser Ser His Tyr Asn Gln Leu Tyr Ser Leu Ser Thr Leu Asp Cys Ser Phe Asn Arg Ile Glu Thr Ser Lys Gly Ile Leu Gln His Phe Pro Lys Ser Leu Ala Phe Phe Asn Leu Thr Asn Asn Ser Val Ala Cys Ile Cys Glu His Gln Lys Phe Leu Gln Trp Val Lys Glu Gln Lys Gln Phe Leu Val Asn Val Glu Gln Met Thr Cys Ala Thr Pro Val Glu Met Asn Thr Ser Leu Val Leu Asp Phe Asn Asn Ser Thr Cys Tyr Met Tyr Lys Thr Ile Ile Ser Val Ser Val Val Ser Val Ile Val Val Ser Thr Val Ala Phe Leu Ile Tyr His Phe Tyr Phe His Leu Ile Leu Ile Ala Gly Cys Lys Lys Tyr Ser Arg Gly Glu Ser Ile Tyr Asp Ala Phe Val Ile Tyr Ser Ser Gln Asn Glu Asp Trp Val Arg Asn Glu Leu Val Lys Asn Leu Glu Glu Gly Val Pro Arg Phe His Leu Cys Leu His Tyr Arg Asp Phe Ile His Gly Val Ala Ile Ala Ala Asn Ile

Ile Gln Glu Gly Phe His Lys Ser Arg Lys Val Ile Val Val Ser 725 730 735

Arg His Phe Ile Gln Ser Arg Trp Cys Ile Phe Glu Tyr Glu Ile Ala 740 745 750

Gln Thr Trp Gln Phe Leu Ser Ser Arg Ser Gly Ile Ile Phe Ile Val 755 760 765

Leu Glu Lys Val Glu Lys Ser Leu Leu Arg Gln Gln Val Glu Leu Tyr 770 775 780

Arg Leu Leu Ser Arg Asn Thr Tyr Leu Glu Trp Glu Asp Asn Pro Leu 785 790 795 800

Gly Arg His Ile Phe Trp Arg Arg Leu Lys Asn Ala Leu Leu Asp Gly 805 810 815

Lys Ala Ser Asn Pro Glu Gln Thr Ala Glu Glu Glu Gln Glu Thr Ala 820 825 830

Thr Trp Thr 835